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## cGAS is required for lethal autoimmune disease in the *Trex1*-deficient mouse model of Aicardi-Goutieres Syndrome<sup>1</sup>

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### Abstract

Detection of intracellular DNA triggers activation of the STING-dependent interferon-stimulatory DNA (ISD) pathway, which is essential for antiviral immune responses. However, chronic activation of this pathway is implicated in autoimmunity. Mutations in *TREX1*, a 3' repair exonuclease that degrades cytosolic DNA, cause Aicardi-Goutieres Syndrome (AGS) and chilblain lupus. *Trex1*<sup>-/-</sup> mice develop lethal, IFN-driven autoimmune disease that is dependent on activation of the ISD pathway, but the DNA sensors that detect the endogenous DNA that accumulates in *Trex1*<sup>-/-</sup> mice have not been defined. Multiple DNA sensors have been proposed to activate the ISD pathway, including cyclic GMP-AMP synthase (cGAS). Here we show that *Trex1*<sup>-/-</sup> mice lacking cGAS are completely protected from lethality, exhibit dramatically reduced tissue inflammation, and fail to develop autoantibodies. These findings implicate cGAS as a key driver of autoimmune disease and suggest that cGAS inhibitors may be useful therapeutics for AGS and related autoimmune diseases.

### Introduction

Detection of intracellular nucleic acids and production of type I IFNs are essential for antiviral immunity. The RNA helicases RIG-I and MDA5 sense intracellular RNA and signal via MAVS to promote IFN production, whereas cGAS detects intracellular DNA ligands and signals via the adaptor STING to trigger the interferon-stimulatory DNA (ISD) response (1). cGAS is a nucleotidyltransferase that directly binds dsDNA in the cytosol and, upon DNA binding, catalyzes the production of cGAMP, which directly binds to and activates STING, triggering production of type I IFNs (2, 3). Studies with cGAS-deficient mice have shown that cGAS is required for the IFN response to transfected DNA ligands, infection with DNA viruses, and retrovirus infection (4-8).

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Although critical for antiviral immunity, innate nucleic acid sensing pathways must be tightly regulated to avoid an IFN response against endogenous nucleic acid ligands. The 3' repair exonuclease Trex1, which degrades cytosolic DNA, is a key regulator of the ISD pathway (9). Loss of function mutations in *TREX1* cause Aicardi-Goutieres syndrome (AGS), a rare autoimmune disease that presents in neonates with clinical features that mimic congenital viral infection (10). AGS patients exhibit elevated type I IFNs, encephalopathy, psychomotor retardation, and premature death (11, 12). Currently, no ameliorative treatments exist for AGS patients (13). *TREX1* mutations also cause familial chilblain lupus (FCL) and retinal vasculopathy with cerebral leukodystrophy (RVCL) and are associated with systemic lupus erythematosus (SLE) (14-18).

*Trex1*<sup>-/-</sup> mice recapitulate many of the key features of AGS and are a tractable model to study AGS disease mechanisms. Trex1-deficient cells accumulate endogenous DNA substrates, including reverse-transcribed retroelement cDNA, that triggers excessive production of type I IFNs resulting in lethal autoimmune disease (9). Autoimmune disease in *Trex1*<sup>-/-</sup> mice is dependent on activation of the ISD pathway, as *Trex1*<sup>-/-</sup> mice that lack STING, IRF3, the type I IFN receptor, or lymphocytes are completely protected from lethality and tissue destruction (9, 19). However, the DNA sensors that detect these endogenous DNA substrates that accumulate in *Trex1*<sup>-/-</sup> mice and trigger IFN production have not been defined. cGAS is a key DNA sensor that activates the ISD response, yet multiple other DNA sensors have been proposed to activate this pathway, including DAI (20), DDX41 (21), and the AIM2-like receptors, IFI16, IFI204, and IFI203 (22-24). One recent study demonstrated that depletion of cGAS within *Trex1*<sup>-/-</sup> mouse embryonic fibroblasts reduced the type I IFN gene signature that is present within these cells (25). However, nothing is known about the *in vivo* role of cGAS in autoimmune pathology. Here, we demonstrate that cGAS is essential for all aspects of the autoimmune disease in *Trex1*<sup>-/-</sup> mice, providing further evidence for its central role as an ISD sensor and revealing a new therapeutic target for AGS and related diseases.

## Materials and Methods

### Mice

We generated *cGAS*<sup>-/-</sup> (*Mb21d1*<sup>-/-</sup>) mice by *in vitro* fertilization of C57BL/6J oocytes with frozen sperm carrying a “knockout first” null allele of *cGAS* from the International Knockout Mouse Consortium (C57BL/6N-Mb21d1<sup>tm1a(EUCOMM)Hmgu</sup>, [http://www.mousephenotype.org/data/alleles/MGI:2442261/tm1a\(EUCOMM\)Hmgu](http://www.mousephenotype.org/data/alleles/MGI:2442261/tm1a(EUCOMM)Hmgu)). Mice were genotyped using the following primers: WT Fwd, GCTGTCTCCACTTGGGCATCT; KO Fwd, AGATGGCGCAACGCAATTAATG; Common Rev, AAGCAGCCACATGAATAGTCTC. *cGAS*-deficient mice with the “knockout first” null allele of *cGAS* were intercrossed with *Trex1*<sup>-/-</sup> mice to generate *cGAS*<sup>-/-</sup>*Trex1*<sup>-/-</sup> mice. *Trex1*<sup>-/-</sup>, *Tmem173*<sup>-/-</sup>, and *Rag2*<sup>-/-</sup> mice have been described (19). All mouse studies were carried out in a specific-pathogen-free facility at the University of Washington with approval of the UW Institutional Animal Care and Use Committee.

## Pathology

All tissues were fixed in 10% neutral buffered formalin and paraffin embedded. Tissue sections (5µm) were stained with hematoxylin and eosin and histological scores were evaluated as described (19).

## Detection of autoantibodies

Heart extracts were prepared and blotted with mouse sera as described (9). Anti-nuclear antibodies were detected with ANA (HEp-2) antigen substrate slides (MBL-BION) with 1:200 diluted mouse sera and anti-mouse IgG Alexa Fluor 488 (Invitrogen) as described (19).

## Cell treatments and analysis

Primary mouse bone marrow-derived macrophages and embryonic fibroblasts were cultured, transfected with nucleic acid ligands, and *Ifnb* mRNA and type I IFN induction were measured as described (26).

## Detection of cGAMP by LC-MS/MS

Mouse hearts were homogenized in extract buffer (50mM HEPES, 2mM EDTA) at a ratio of 0.1g heart tissue per 1ml of buffer. Ice-cold methanol was added to 80% final concentration. Extracts were spiked with isotope-labeled cGAMP (cGAMP\*) at 10nM, sonicated, cleared three times by centrifugation at  $16,000 \times g$ , lyophilized, and resuspended in water. Detection of cGAMP in heart lysates was performed with LC-MS/MS as described (27) with the following modifications. Isotope-labeled cGAMP (cGAMP\*, 12C1013C10H2415N514N5O13P2, MW=689 g/mol), synthesized with recombinant cGAS and per labeled  $^{13}C$ ,  $^{15}N$ -ATP and natural abundance GTP, was used as an internal standard. Both cGAMP and cGAMP\* were eluted at 2.7 min. The following MRM transitions were detected: cGAMP: +675.1/152.1 (qualifier), +675.1/136.0 (quantifier), and cGAMP\*: +690.0/146.0 (quantifier). A standard curve was constructed using standard 1 (250 nM cGAMP\*, 10 nM cGAMP) and standard 2 (250 nM cGAMP\*, 1000 nM cGAMP).

## Results and Discussion

### cGAS is required for lethal autoimmunity in *Trex1*-deficient mice

To explore cGAS function *in vivo*, we generated cGAS-deficient mice from frozen sperm carrying a null allele of cGAS (*cGAS*<sup>-/-</sup>, also referred to as *Mb21d1*<sup>-/-</sup>, **Supplementary Figure 1A,B**). cGAS-deficient mice were physically indistinguishable from their wild type littermates (data not shown). Moreover, consistent with recent studies (5), the IFN response to transfected DNA in *cGAS*<sup>-/-</sup> bone marrow macrophages (**Supplementary Figure 2A,B**) and embryonic fibroblasts (**Supplementary Figure 2C,D**) was dramatically impaired and similar to STING-deficient cells, while the response to transfected RNA remained intact.

To determine whether cGAS is required for autoimmune disease in *Trex1*-deficient mice, we intercrossed *Trex1*<sup>-/-</sup> and *cGAS*<sup>-/-</sup> mice and monitored survival. *Trex1* and *cGAS* are genetically linked, separated by 16 cM on chromosome 9. For this reason, recovery of *cGAS*<sup>-/-</sup>*Trex1*<sup>-/-</sup> mice from intercrosses of *cGAS*<sup>+/-</sup>*Trex1*<sup>+/-</sup> mice occurred at a much lower

frequency than would be expected with unlinked genes. In contrast to  $cGAS^{+/+}Trex1^{-/-}$  mice, which succumbed to lethal disease as expected,  $cGAS^{-/-}Trex1^{-/-}$  mice were completely protected from lethality (**Figure 1**).  $Trex1^{-/-}$  mice heterozygous for  $cGAS$  were partially rescued from death similar to  $Irf3^{+/-}Trex1^{-/-}$  mice (9). This contrasts with  $Trex1^{-/-}$  mice lacking a single copy of *Tmem173* (which encodes STING), which are completely protected from lethal disease (19).

### **cGAS promotes autoimmune tissue destruction in $Trex1^{-/-}$ mice**

$Trex1$ -deficient mice develop multiorgan inflammation, including severe inflammatory myocarditis that is dependent on type I IFNs and STING (9, 19). To determine whether  $cGAS$  is required to drive autoimmune tissue destruction, we performed a blinded histological analysis of tissues from  $cGAS^{-/-}Trex1^{-/-}$  mice compared to  $cGAS^{+/+}Trex1^{-/-}$ ,  $cGAS^{+/-}Trex1^{-/-}$  and  $cGAS^{-/-}Trex1^{+/+}$  mice. Consistent with previous studies, we found evidence of inflammation in multiple tissues from  $cGAS^{+/+}Trex1^{-/-}$  mice, including the heart, skin, glandular stomach, skeletal muscle, tongue, and kidney, whereas the brain was unaffected (**Figure 2**). Heart tissue from  $Trex1^{-/-}$  mice was characterized by focally extensive subendocardial and perivascular lymphoplasmacytic myocarditis (**Figure 2A**, left panel). Skin tissue exhibited focally extensive proliferative dermatitis with lymphoplasmacytic and lesser neutrophilic cellulitis and myositis (**Figure 2A**, middle panel). Finally, the glandular stomach was characterized by diffuse moderate proliferative gastritis with multifocal lymphoid aggregates (**Figure 2A**, right panel). In contrast, analysis of heart and skin tissue from  $cGAS^{-/-}Trex1^{-/-}$  revealed no signs of inflammation (**Figure 2A,B**). Mild inflammation was observed in glandular stomach, skeletal muscle, tongue, and kidney tissue from  $cGAS^{-/-}Trex1^{-/-}$  mice, which was similar to  $cGAS^{-/-}Trex1^{+/+}$  mice and clearly distinct from that observed in  $cGAS^{+/+}Trex1^{-/-}$  mice (**Figure 2B** and data not shown). For example, multifocal lymphoid aggregates were not detected in glandular stomach tissue from  $cGAS^{-/-}Trex1^{-/-}$  mice and the mild inflammation observed in these mice was in a distinct region of the stomach (*margo plicatus*/limiting ridge) compared to  $cGAS^{+/+}Trex1^{-/-}$  mice (**Fig. 2A** and data not shown). Thus,  $cGAS^{-/-}Trex1^{-/-}$  mice are protected from tissue destruction, similar to  $Tmem173^{-/-}Trex1^{-/-}$  mice (9, 19), indicating that the  $cGAS$ /STING ISD pathway is required to drive autoimmune tissue destruction in  $Trex1$ -deficient mice.

### **cGAS drives the autoantibody response in $Trex1^{-/-}$ mice**

$Trex1$ -deficient mice develop a T cell-dependent autoantibody response against heart tissue antigens that is dependent on IRF3 and type I IFN signaling (9, 19). To determine whether  $cGAS$  is required for the development of these autoantibodies, we blotted heart tissue extracts with sera from  $cGAS^{+/+}Trex1^{-/-}$ ,  $cGAS^{+/-}Trex1^{-/-}$ , and  $cGAS^{-/-}Trex1^{-/-}$  mice. Sera from multiple  $cGAS^{+/+}Trex1^{-/-}$  and  $cGAS^{+/-}Trex1^{-/-}$  mice exhibited strong autoreactivity against heart extracts (**Figure 3A**, left panels). The antigen specificities of these heart autoantibodies were variable among different mice, consistent with our previous analysis of  $Trex1$ -deficient mice (9, 19). In contrast, minimal autoreactivity, similar to that observed in  $Trex1^{+/-}$  and  $Trex1^{+/+}$  mice, was observed in sera from  $cGAS^{-/-}Trex1^{-/-}$  mice (**Figure 3A**,

right panels). This demonstrates that cGAS is required for the development of autoantibodies against heart tissue antigens in *Trex1*-deficient mice.

In addition to the heart-specific autoantibody response, *Trex1*-deficient mice develop several hallmarks of lupus including anti-nuclear antibodies (ANAs) (19). Whether the development of ANAs in *Trex1*-deficient mice requires the cGAS/STING ISD pathway has not yet been tested. To explore this question, we evaluated sera from *cGAS*<sup>+/+</sup>*Trex1*<sup>-/-</sup>, *cGAS*<sup>+/-</sup>*Trex1*<sup>-/-</sup>, and *cGAS*<sup>-/-</sup>*Trex1*<sup>-/-</sup> mice for the presence of ANAs by staining HEP-2 cell-coated slides. Consistent with previous work, ANAs with variable signal intensity were observed in sera from all *cGAS*<sup>+/+</sup>*Trex1*<sup>-/-</sup> mice (n=7/7) and the majority of *cGAS*<sup>+/-</sup>*Trex1*<sup>-/-</sup> mice (n=6/7) (**Figure 3B**). In contrast, ANAs were not detected in sera from age-matched *cGAS*<sup>-/-</sup>*Trex1*<sup>-/-</sup> mice (n=0/3) (**Figure 3B**) indicating that cGAS is required for the development of ANAs in *Trex1*-deficient mice.

### cGAS promotes type I IFN production in *Trex1*<sup>-/-</sup> mice

Disease in *Trex1*-deficient mice is driven by type I IFN signaling, as *Ifnar*<sup>-/-</sup>*Trex1*<sup>-/-</sup> mice are protected from disease (9). To determine whether cGAS promotes type I IFN production in *Trex1*<sup>-/-</sup> mice, we measured a panel of IFN-stimulated genes (ISGs) in total peripheral blood cells from *cGAS*<sup>+/+</sup>*Trex1*<sup>-/-</sup>, *cGAS*<sup>-/-</sup>*Trex1*<sup>-/-</sup> and *Trex1*<sup>+/+</sup> mice. We detected a significant increase in expression of two ISGs, *Cxcl10* and *Mx1*, and a trend towards increased expression of *Isg15* and *Ifit2* in cells from *cGAS*<sup>+/+</sup>*Trex1*<sup>-/-</sup> mice compared to *Trex1*<sup>+/+</sup> mice, while ISG expression was similar in blood cells from *cGAS*<sup>-/-</sup>*Trex1*<sup>-/-</sup> and *Trex1*<sup>+/+</sup> mice (**Figure 4A**).

In a prior study, we identified a localized focus of IFN-responsive cells at the endocardial surface of the heart in neonatal *Trex1*-deficient mice, suggesting that disease is initiated in this tissue. To explore the possibility that cGAS is activated in *Trex1*<sup>-/-</sup> hearts, we used mass spectrometry to measure cGAMP in *Trex1*<sup>-/-</sup> heart tissue. We reproducibly detected elevated levels of cGAMP in heart tissue extracts from *cGAS*<sup>+/+</sup>*Trex1*<sup>-/-</sup> mice, while the cGAMP was low or undetectable in extracts from *cGAS*<sup>-/-</sup>*Trex1*<sup>-/-</sup> and *Trex1*<sup>+/+</sup> mice (**Figure 4B**). We hypothesize that cGAS activation in heart tissue drives lethal myocarditis in *Trex1*<sup>-/-</sup> mice. *Trex1*-deficient mice with a conditional allele of cGAS (**Supplementary Fig. 1A**) will allow for selective depletion of cGAS in future experiments aiming to define the cell types in which cGAS is required to drive disease.

We provide clear genetic evidence that cGAS is absolutely required for disease in *Trex1*-deficient mice. *cGAS*<sup>-/-</sup>*Trex1*<sup>-/-</sup> mice are completely protected from lethality, lack autoimmune inflammation, fail to develop autoantibodies, and lack the peripheral blood IFN signature that is present in plain *Trex1*<sup>-/-</sup> mice. Moreover, we show that the inflamed hearts of *Trex1*<sup>-/-</sup> mice contain elevated levels of cGAMP, thus demonstrating *in vivo* activation of cGAS during autoimmune inflammation. While we cannot rule out the possibility that other DNA sensors proposed to activate the STING-dependent IFN response may contribute to autoimmunity in *Trex1*-deficient mice, our data suggest that, if other sensors participate, they must do so by functioning together with cGAS, and not as independent sensors. Altogether, our studies demonstrate that cGAS is a key driver of autoimmune disease and

suggest that cGAS inhibitors may be useful therapeutics for AGS patients with *TREX1* mutations.

## Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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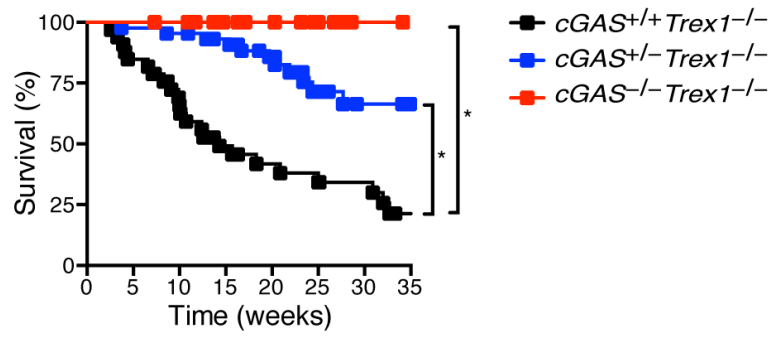
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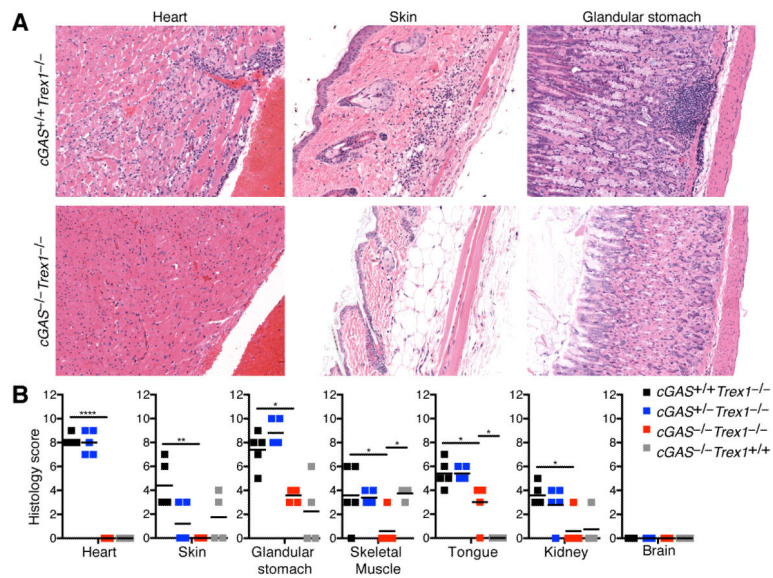
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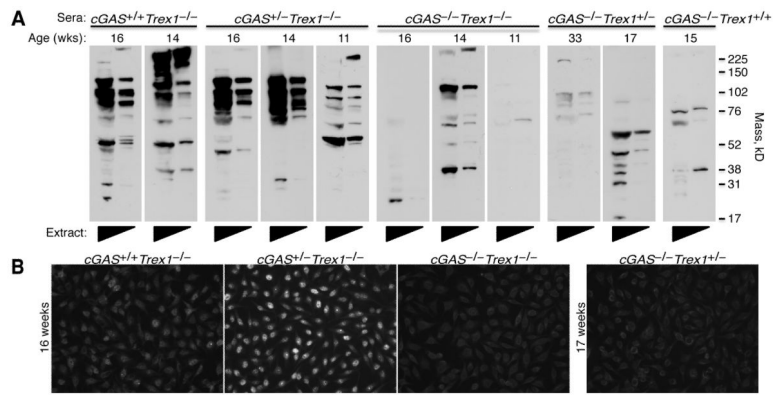
**Figure 1. cGAS is required for lethal autoimmune disease in *Trex1*-deficient mice**

Survival curves of *cGAS*<sup>+/+</sup>*Trex1*<sup>-/-</sup> (n=33), *cGAS*<sup>+/-</sup>*Trex1*<sup>-/-</sup> (n=44), and *cGAS*<sup>-/-</sup>*Trex1*<sup>-/-</sup> (n=29) mice. Both *cGAS*-deficient and *Trex1*-deficient mice were on a pure C57BL/6 background. Statistical analysis was performed with a Log-rank (Mantel-Cox) test. \*p < 0.0001.



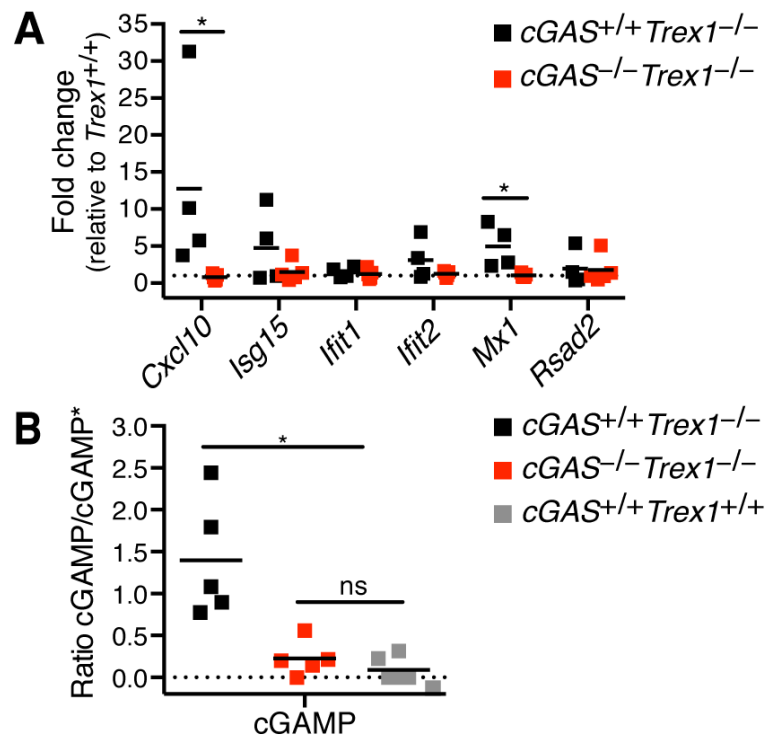
**Figure 2. cGAS drives tissue inflammation in Trex1-deficient mice**

(A) Representative H&E-stained heart, skin, and glandular stomach tissue sections from  $cGAS^{+/+}Trex1^{-/-}$  and  $cGAS^{-/-}Trex1^{-/-}$  mice (20X original magnification). (B) Histological scores of inflammation in the indicated tissues from  $cGAS^{+/+}Trex1^{-/-}$ ,  $cGAS^{+/-}Trex1^{-/-}$ ,  $cGAS^{-/-}Trex1^{-/-}$  and  $cGAS^{-/-}Trex1^{+/+}$  mice. All histological analysis was performed in a blinded manner. Data are representative of at least four mice of each genotype. Statistical analysis was performed comparing  $cGAS^{-/-}Trex1^{-/-}$  to  $cGAS^{+/+}Trex1^{-/-}$  and  $cGAS^{-/-}Trex1^{+/+}$  mice using a one-way ANOVA with Tukey's multiple comparison posttest. \* $p < 0.05$ , \*\* $p < 0.005$ , \*\*\* $p < 0.0001$ .



### Figure 3. cGAS-dependent autoantibodies in *Trex1*-deficient mice

(A) Autoantibodies against heart antigens evaluated by blotting neat and 1:5 diluted heart extracts from *Rag2*<sup>-/-</sup>*Trex1*<sup>-/-</sup> mice with sera from mice of the indicated genotype. Immunoblots were prepared with sera from different mice harvested at the age indicated, rather than with sera harvested repeatedly from a single mouse. Data are representative of two independent experiments. (B) Antinuclear antibodies (ANA) in sera from littermate *cGAS*<sup>+/+</sup>*Trex1*<sup>-/-</sup>, *cGAS*<sup>+/-</sup>*Trex1*<sup>-/-</sup>, and *cGAS*<sup>-/-</sup>*Trex1*<sup>-/-</sup> mice (left panels) analyzed by immunofluorescence staining of HEp-2 cell-coated slides. ANA in serum from *cGAS*<sup>-/-</sup>*Trex1*<sup>+/-</sup> mice was analyzed as a control (right panel). Data are representative of three independent experiments with a total of at least three mice of each genotype.



**Figure 4. cGAS promotes type I IFN production in *Trex1*-deficient mice**

(A) Quantification by RT-PCR of a panel of IFN-stimulated gene transcripts in total peripheral blood cells from *cGAS*<sup>+/+</sup>*Trex1*<sup>-/-</sup> and *cGAS*<sup>-/-</sup>*Trex1*<sup>-/-</sup> mice relative *Trex1*<sup>+/+</sup> mice. Data are representative of one independent experiment with at least four mice of each genotype. (B) Analysis of cGAMP production in whole heart extracts from *cGAS*<sup>+/+</sup>*Trex1*<sup>-/-</sup>, *cGAS*<sup>-/-</sup>*Trex1*<sup>-/-</sup>, and *Trex1*<sup>+/+</sup> mice by LC-MS/MS. Data are plotted relative to the internal standard (cGAMP\*). Data are representative of one independent experiment with at least five mice of each genotype. Statistical analysis was performed with a Mann-Whitney test. \**p* < 0.05.